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(54) Title: MONOCLONAL ANBIBODIES WHICH BIND TO TUMOR REJECTION ANTIGEN PRECURSOR MAGE-1, RECOMBI-NANT MAGE-1, AND MAGE-1 DERIVED IMMUNOGENIC PEPTIDES

(57) Abstract

The invention relates to monoclonal antibodies which specifically bind to the tumor rejection antigen precursor molecule MAGE-1, hybridomas which produce these monoclonal antibodies, and their use. Also described is a recombinant form of MAGE-1, peptides which are useful as immunogens, and immunogenic compositions containing the peptides and an adjuvant.

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7

MONOCLONAL ANTIBODIES WHICH BIND TO TUMOR REJECTION
ANTIGEN PRECURSOR MAGE-1, RECOMBINANT MAGE -1,
AND MAGE-1 DERIVED IMMUNOGENIC PEPTIDES

RELATED APPLICATION

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This application is a continuation-in-part of Serial No. 037,230 filed March 26, 1993, which is itself a continuation-in-part of PCT Application PCT/US92/04354 filed on May 22, 1992 designating the United States, which is a continuation-in-part of Serial Number 807,043, filed December 12, 1991, which is a continuation-in-part of Serial Number 764,365, filed September 23, 1991, which is a continuation-in-part of Serial Number 728,838, filed July 9, 1991, which is a continuation-in-part of Serial Number 705,702, filed May 23, 1991, and now abandoned.

FIELD OF THE INVENTION

This invention relates in general to the field of immunogenetics as applied to the study of oncology. More specifically, it relates to the study and analysis of mechanisms by which tumors are recognized by the organism's immune system such as through the presentation of so-called tumor rejection antigens, and the expression of what will be referred to herein as "tumor rejection antigen precursors" or "TRAPs". Most specifically, it refers to one such TRAP, i.e., MAGE-1, produced recombinantly, and monoclonal antibodies and antisera directed against MAGE-1, as well as their use.

BACKGROUND AND PRIOR ART

The study of the recognition or lack of recognition of cancer cells by a host organism has proceeded in many different directions. Understanding of the field presumes some understanding of both basic immunology and oncology.

Early research on mouse tumors revealed that these displayed molecules which led to rejection of tumor cells when transplanted into syngeneic animals. These molecules are "recognized" by T-cells in the recipient animal, and provoke a cytolytic T-cell response with lysis of the transplanted cells. This evidence was first obtained with tumors induced in vitro by chemical carcinogens, such as methylcholanthrene.

2

The antigens expressed by the tumors and which elicited the Tcell response were found to be different for each tumor. See Prehn, et al., J. Natl. Canc. Inst. 18: 769-778 (1957); Klein et al., Cancer Res. 20: 1561-1572 (1960); Gross, Cancer Res. 3: 326-333 (1943), Basombrio, Cancer Res. 30: 2458-2462 (1970) for general teachings on inducing tumors with chemical carcinogens and differences in cell surface antigens. class of antigens has come to be known as "tumor specific transplantation antigens" or "TSTAs". Following observation of the presentation of such antigens when induced by chemical carcinogens, similar results were obtained when tumors were induced in vitro via ultraviolet radiation. Kripke, J. Natl. Canc. Inst. 53: 333-1336 (1974).

While T-cell mediated immune responses were observed for the types of tumor described <u>supra</u>, spontaneous tumors were thought to be generally non-immunogenic. These were therefore believed not to present antigens which provoked a response to the tumor in the tumor carrying subject. See Hewitt, et al., Brit. J. Cancer 33: 241-259 (1976).

The family of tum antigen presenting cell lines are immunogenic variants obtained by mutagenesis of mouse tumor cells or cell lines, as described by Boon et al., J. Exp. Med. 152: 1184-1193 (1980), the disclosure of which is incorporated by reference. To elaborate, tum antigens are obtained by mutating tumor cells which do not generate an immune response in syngeneic mice and will form tumors (i.e., "tum*" cells). When these tum* cells are mutagenized, they are rejected by syngeneic mice, and fail to form tumors (thus "tum"). See Boon et al., Proc. Natl. Acad. Sci. USA 74: 272 (1977), the disclosure of which is incorporated by reference. Many tumor types have been shown to exhibit this phenomenon. See, e.g., Frost et al., Cancer Res. 43: 125 (1983).

It appears that tum variants fail to form progressive tumors because they elicit an immune rejection process. The evidence in favor of this hypothesis includes the ability of "tum" variants of tumors, i.e., those which do not normally form tumors, to do so in mice with immune systems suppressed

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by sublethal irradiation, Van Pel et al., Proc. Natl, Acad. Sci. USA 76: 5282-5285 (1979); and the observation that intraperitoneally injected tum cells of mastocytoma P815 multiply exponentially for 12-15 days, and then are eliminated in only a few days in the midst of an influx of lymphocytes and macrophages (Uyttenhove et al., J. Exp. Med. 152: 1175-1183 (1980)). Further evidence includes the observation that mice acquire an immune memory which permits them to resist subsequent challenge to the same tum variant, even when immunosuppressive amounts of radiation are administered with the following challenge of cells (Boon et al., Proc. Natl, Acad. Sci. USA 74: 272-275 (1977); Van Pel et al., supra; Uyttenhove et al., supra). Later research found that when spontaneous tumors were subjected to mutagenesis, immunogenic variants were produced which did generate a response. these variants were able to elicit an immune protective response against the original tumor. See Van Pel et al., J. Exp. Med. 157: 1992-2001 (1983). Thus, it has been shown that it is possible to elicit presentation of a so-called "tumor rejection antigen" in a tumor which is a target for a syngeneic rejection response. Similar results have been obtained when foreign genes have been transfected into spontaneous tumors. See Fearson et al., Cancer Res. 48: 2975-1980 (1988) in this regard.

A class of antigens has been recognized which are presented on the surface of tumor cells and are recognized by cytotoxic T cells, leading to lysis. This class of antigens will be referred to as "tumor rejection antigens" or "TRAs" hereafter. TRAs may or may not elicit antibody responses. The extent to which these antigens have been studied, has been via cytolytic T cell characterization studies in vitro i.e., the study of the identification of the antigen by a particular cytolytic T cell ("CTL" hereafter) subset. The subset proliferates upon recognition of the presented tumor rejection antigen, and the cells presenting the antigen are lysed. Characterization studies have identified CTL clones which specifically lyse cells expressing the antigens. Examples of

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this work may be found in Levy et al., Adv. Cancer Res. 24: 1-59 (1977); Boon et al., J. Exp. Med. 152: 1184-1193 (1980); Brunner et al., J. Immunol. 124: 1627-1634 (1980); Maryanski et al., Eur. J. Immunol. 124: 1627-1634 (1980); Maryanski et al., Eur. J. Immunol. 12: 406-412 (1982); Palladino et al., Canc. Res. 47: 5074-5079 (1987). This type of analysis is required for other types of antigens recognized by CTLs, including major histocompatibility antigens, the male specific H-Y antigens, and a class of antigens, referred to as "tum-" antigens, and discussed herein.

A tumor exemplary of the subject matter described supra is known as P815. See DePlaen et al., Proc. Natl. Acad. Sci. USA 85: 2274-2278 (1988); Szikora et al., EMBO J 9: 1041-1050 (1990), and Sibille et al., J. Exp. Med. 172: 35-45 (1990), the disclosures of which are incorporated by reference. P815 tumor is a mastocytoma, induced in a DBA/2 mouse with methylcholanthrene and cultured as both an in vitro tumor and a cell line. The P815 line has generated many tum variants following mutagenesis, including variants referred to as P91A (DePlaen, supra), 35B (Szikora, supra), and P198 (Sibille, supra). In contrast to tumor rejection antigens - and this is a key distinction - the tum antigens are only present after the tumor cells are mutagenized. Tumor rejection antigens are present on cells of a given tumor without mutagenesis. Hence, with reference to the literature, a cell line can be tum', such as the line referred to as "P1", and can be provoked to produce tum variants. Since the tum phenotype differs from that of the parent cell line, one expects a difference in the DNA of tum cell lines as compared to their tum parental lines, and this difference can be exploited to locate the gene of interest in tum cells. As a result, it was found that genes of tum variants such as P91A, 35B and P198 differ from their normal alleles by point mutations in the coding regions of the gene. See Szikora and Sibille, supra, and Lurquin et al., Cell 58: 293-303 (1989). This has proved <u>not</u> to be the case with the TRAs of this invention. These papers also demonstrated that peptides derived from the tum antigen are

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presented by the L^d molecule for recognition by CTLs. P91A is presented by L^d , P35 by D^d and P198 by K^d .

Prior patent applications PCT/US92/04354, U.S. Serial No. 807,043; 764,364; 728,838 and 707,702, all of which are incorporated by reference, describe inventions involving, inter alia, genes and other nucleic acid molecules which code for various TRAPs, which are in turn processed to tumor rejection antigen, or "TRAS".

The genes are useful as a source for the isolated and purified tumor rejection antigen precursor and the TRA themselves, either of which can be used as an agent for treating the cancer for which the antigen is a "marker", as well as in various diagnostic and surveillance approaches to oncology, discussed infra. It is known, for example, that tum cells can be used to generate CTLs which lyse cells presenting different tum antigens as well as tum cells. See, e.g., Maryanski et al., Eur. J. Immunol 12: 401 (1982); and Van den Eynde et al., Modern Trends in Leukemia IX (June 1990), the disclosures of which are incorporated by reference. The tumor rejection antigen precursor may be expressed in cells transfected by the gene, and then used to generate an immune response against a tumor of interest.

In the parallel case of human neoplasms, it has been observed that autologous mixed lymphocyte-tumor cell cultures ("MLTC" hereafter) frequently generate responder lymphocytes which lyse autologous tumor cells and do not lyse natural killer targets, autologous EBV-transformed B cells, autologous fibroblasts (see Anichini et al., Immunol. Today 8: 385-389 (1987)). This response has been particularly well studied for melanomas, and MLTC have been carried out either with peripheral blood cells or with tumor infiltrating lymphocytes. Examples of the literature in this area including Knuth et al., Proc. Natl. Acad. Sci. USA 86: 2804-2802 (1984); Mukherji et al., J. Exp. Med. 158: 240 (1983); Hérin et all, Int. J. Canc. 39: 390-396 (1987); Topalian et al, J. Clin. Oncol 6: 839-853 (1988). Stable cytotoxic T cell clones ("CTLs" hereafter) have been derived from MLTC

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responder cells, and these clones are specific for the tumor cells. See Mukherji et al., supra, Hérin et all, supra, Knuth et al., supra. The antigens recognized on tumor cells by these autologous CTLs do not appear to represent a cultural artifact, since they are found on tumor cells in vivo. Topalian et al., supra; Degiovanni et al., Eur. J. Immunol. 20: 1865-1868 (1990). These observations, coupled with the techniques used herein to isolate the genes for specific murine tumor rejection antigen precursors, have led to the isolation of nucleic acid sequences coding for tumor rejection antigen precursors of TRAs presented on human tumors. now possible to isolate the nucleic acid sequences which code for tumor rejection antigen precursors, including, but not being limited to those most characteristic of a particular tumor, with ramifications that are described infra.

Additional work has focused upon the presentation of TRAs by the class of molecules known as human leukocyte antigens, This work has resulted in several unexpected discoveries regarding the field. Specifically in U.S. patent application Serial Number 938,334, the disclosure of which is incorporated by reference, nonapeptides are taught which are presented by the HLA-A1 molecule. The reference teaches that given the known specificity of particular peptides for particular HLA molecules, one should expect a particular peptide to bind one HLA molecule, but not others. important, because different individuals possess different HLA phenotypes. As a result, while identification of a particular peptide as being a partner for a specific HLA molecule has diagnostic and therapeutic ramifications, these are only relevant for individuals with that particular HLA phenotype. There is a need for further work in the area, because cellular abnormalities are not restricted to one particular HLA phenotype, and targeted therapy requires some knowledge of the phenotype of the abnormal cells at issue.

In U.S. Patent Application Serial Number 008,446, filed January 22, 1993 and incorporated by reference, the fact that the MAGE-1 expression product is processed to a second TRA is

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disclosed. This second TRA is presented by HLA-C-clone-10 molecules. The disclosure shows that a given TRAP can yield a plurality of TRAs.

In U.S. Patent Application Serial Number 994,928, filed December 22, 1992, and incorporated by reference herein, tyrosinase is described as a tumor rejection antigen precursor. This reference discloses that a molecule which is produced by some normal cells (e.g., melanocytes), is processed in tumor cells to yield a tumor rejection antigen that is presented by HLA-A2 molecules.

The prior applications cited supra discussed antibodies against tumor rejection antigen precursors generally. The present investigators have utilized the isolated nucleic acid molecule coding for MAGE-1 to produce a recombinant MAGE-1 protein, and peptides derived therefrom. These have been used to produce polyclonal and monoclonal antibodies which specifically bind to MAGE-1. These antibodies, and their use, constitute the invention described and claimed herein.

BRIEF DESCRIPTION OF THE FIGURES

Figure 1 shows, schematically, the MAGE-1 gene, oligopeptides derived from the recombinant MAGE-1 protein, and comparison with corresponding sequences in MAGE-2 and MAGE-3 deduced amino acid sequences.

Figure 2A shows silver stained SDS-polyacrylamide gels of affinity purified, MAGE-1 recombinant protein. Figure 2B presents immunoblotting work where recombinant MAGE-1 protein was used against rabbit antisera derived from immunization with three peptides (SEQ ID NOS: 2, 3 and 4). Blotting was at 1:1000 dilution. As a control, recombinant mouse p53 was used.

Figure 3A shows the reactivity pattern of mAb MA 454 against six melanoma lines. Figure 3B shows the results obtained using rabbit polyclonal antisera against the same lines. Figure 3C shows results obtained with a MAGE-1 transfected cell line (MZ2-MEL 2.2-ET.1), and its parent (MZ2-MEL 2.2).

Figure 4 presents immunoblot analysis using the

antibodies against tissue lysates.

DETAILED DESCRIPTION OF PREFERRED EMBODIMENTS

Many different "MAGE" genes have been identified, as will be seen from the applications and references cited supra. The MAGE-1 gene is at issue on the present case, and is the only one discussed hereafter. For convenience, it is presented herein as SEQ ID NO: 1.

"MAGE" as used herein refers to a nucleic acid sequence isolated from human cells.

When "TRAP" or "TRAS" are discussed herein as being specific to a tumor type, this means that the molecule under consideration is associated with that type of tumor, although not necessarily to the exclusion of other tumor types.

Example 1

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The cell line MZ2-MEL 3.1 described in, e.g., Van den Eynde et al., Int. J. Cancer 44: 634-640 (1989) and in the parent application cited <u>supra</u>, previously observed to express MAGE-1, was used as a source of total RNA. The total RNA was extracted from the cells, and was then subjected to reverse transcription/polymerase chain reaction, using the primers CHO8 and CHO9, as described by Van der Bruggen et al., Science 254: 1643-1647 (Dec. 13, 1991), the disclosure of which is incorporated by reference. This paper describes the "RT-PCR" technique, as does the Brasseur et al., Int. J. Cancer 52: 839-841 (1992). It must be understood, however, that the sequence of MAGE-1 is known to the art, and other primers could be used besides CHO8 and CHO9.

Once the RT-PCR protocols were completed, the products were cloned directly into plasmid pT7 Blue (Novagen, Madison WI), following manufacturer's instructions which constituted well known techniques. Following the cloning, the recombinant plasmid DNA was treated with restriction endonucleases to generate fragments which included fragments containing the MAGE-1 gene. See, e.g. Van der Bruggen et al, <u>supra</u>.

The appropriate cDNA insert was subcloned unidirectionally, into plasmids pQE9, pQE10 and pQE11, using BamHI and HindIII cloning sites in pT7 Blue. The plasmids

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were transfected into \underline{E} . $\underline{\operatorname{coli}}$, and recombinant protein production was induced via IPTG, as the host plasmid contains the lac operon. This yielded a fusion protein containing the MAGE-1 polypeptide sequence, which could be purified via Ni^{2+} ion affinity chromatography.

The DNA sequence of the recombinant clone was obtained, and was confirmed to encode 163 amino acids which correspond to deduced amino acids 57-219 of predicted MAGE-1 amino sequence, plus 30 residues from the plasmid itself. Figure 1 shows this. The expected molecular mass is about 20-22 kDa.

When clones in pQE10 were studied, indeed, a recombinant protein of about 20 kDa was produced following IPTG induction. Other minor protein species of 70 kDa, 43 kDa, 17 kDa and 15 kDa were also found, as is seen in figure 2A.

Example 2

The following describes procedures used to produce antibodies to MAGE-1. Based upon the predicted MAGE-1 amino acid sequence, three oligopeptides were prepared:

Ile Asn Phe Thr Arg Gln Arg Gln Pro Ser Glu Gly Ser Ser (SEQ ID NO: 2)

Leu Phe Arg Ala Val Ile Thr Lys Lys Val Ala Asp
(SEQ ID No: 3)

Asp Val Lys Glu Ala Asp Pro Thr Gly His Ser Tyr
(SEQ ID No: 4)

Rabbits were immunized with the peptides, and were then treated to collect antiserum.

Antisera prepared against these three peptides were then used with \underline{E} . $\underline{\operatorname{coli}}$ produced, recombinant MAGE-1 protein, in immunoblotting experiments. The results, set forth in figure 2B, show that only antiserum raised against the first of these peptides, i.e., SEQ ID NO: 2 reacted strongly. The fact that additional protein species that copurified with the 20 kDa fusion protein also showed reactivity, suggests that these are

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aggregates of the fusion protein. The peptide used corresponds to deduced amino acids 68-81 of the MAGE-1 of the predicted MAGE-1 protein.

When immunoblotting was carried out using lysates of melanoma cell line MZ2-MEL 3.1, no detectable MAGE protein was found.

Example 3

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Monoclonal antibodies were then prepared. Purified recombinant protein, produced as described supra, was used to immunize BALB/C mice. Hybridomas were generated and cloned. The protocol used was that described by Dippold et al., Proc. Natl. Acad. Sci. USA 77: 6114-6118 (1980), the disclosure of which is incorporated by reference. The key difference, of course, was the immunogen used for immunization.

Once hybridomas were generated, their supernatants were screened using a standard, solid phase ELISA on microtiter plates, using the immunizing fusion protein as target antigen. Five clones were found to be reactive. They all also showed moderate to strong reactivity in immunoblots.

As a control, mouse p53 protein, expressed in the same plasmid vector, was also tested. No reactivity was seen. These results are summarized in Table 1 which follows:

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TABLE 1. Reactivity of mouse anti-recombinant MAGE-1, mabs toward recombinant MAGE-1 protein and control p53 protein

Assay	ELISA	A	Immunoblot		
mAb	MAGE-1	p 53	MAGE-1	p53	
MA32	++*	-	++#	_	
MA231	+	-	++	_	
MA399	++	-	++	-	
MA430	++	_	+++	_	
MA454	++	_	+++	-	

*ELISA titer using hybridoma supernatants: -, <1:16; +, 1:64; ++, 1:256. #Immunoblot signal intensity: -, negative; +, weak; ++, moderate; +++, strong.

Example 4

The mAbs described <u>supra</u> were then tested against lysates of melanoma cell lines. The cell lines tested, i.e., MZ2-MEL 3.1, MZ2-MEL 2.2, and SK-MEL-187, are all well known. MZ2-MEL 2.2 is a MAGE-1 loss variant derived from MAGE-1 positive parental MZ2-MEL 3.1 by CTL selection (van der Bruggen et al., Int. J. Cancer 44: 634-640 (1989)). These cells had been "typed" by RT-PCR as being MAGE-1*2*3*(MZ2-MEL 3.1), MAGE-1*2*3*(MZ2-MEL 2.2), and MAGE 1*2*3*(SK MEL-187). The lysates were prepared by homogenizing the cells in Nonidet P40 (NP-40) buffer (1% NP-40, 50 mM Tris-HCl, pH 8.0, 150 mM NaCl). The results are shown in figure 3A.

Monoclonal antibody MA 454 reacted with a 46 kDa protein present in MZ2-MEL 3.1 lysate, but not in lysates of either of the other two cell lines. When three additional melanoma

12

lines were tested, only those which were typed as being MAGE-1 positive reacted with the mAb. Expression of MAGE-2 or MAGE-3 was irrelevant.

The polyclonal antiserum described supra, was also tested against these lysates. Results are shown in figure 3B. It was positive for MZ2-MEL 3.1, and for MAGE-1 transfected cell line MZ2-MEL 2.2-ET.1, but was negative for parental line MZ2-MEL 2.2.

Example 5

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Lysates were prepared from liver, kidney and testis tissue, and from four melanoma cell lines including one MAGE-1*2*3* line, two MAGE-1*2*3*, and one MAGE-1*2*3* lines. The lysates were prepared as described supra. When immunoblotting was carried out, testis lysates were positive with mAb 454, as were MAGE-1 positive melanomas. No other lysates were positive which is in complete agreement with mRNA expression data.

The same experiments were carried out using polyclonal antiserum, and the results paralleled those for the monoclonal antibodies. Figure 4 presents these results.

The foregoing experiments describe the production of monoclonal antibodies which specifically bind to a tumor rejection antigen precursor TRAP. The studies show binding both to the "wild type" MAGE-1 molecule, and the recombinant form, but not to either of MAGE-2 or MAGE-3. A particularly preferred species of MAGE-1 binding mAb, i.e., MA454, has been deposited at the American Type Culture Collection under Accession Number HB 11540.

The invention thus relates to MAGE-1 specific monoclonal antibodies and the hybridomas which produce them. The mAbs were found to be useful in determining expression of MAGE-1 in cell lysates. Specifically, the mAbs can be added, e.g., in labelled form, bound to a solid phase, or otherwise treated to increase the sensitivity of MAGE-1 detection. Any of the standard types of immunoassays, including ELISAs, RIAs, competitive assays, agglutination assays, and all others are encompassed with respect to the way the mAbs can be used.

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"Cell lysate" as used herein refers not only to a sample which is expressly lysed, but also to those samples which contain cells which have been lysed in vivo, or any sample which contains material normally internal to the cells. The detection of MAGE-1 expression product is useful, e.g., in diagnosing or monitoring the presence or progress of a cancer.

The isolated, recombinant MAGE-1 protein is also a feature of this invention. This molecule has a molecular weight of about 20-22 kDa as determined by SDS-PAGE, and is useful as an immunogen as are the peptides of SEQ ID NOS: 2, 3 and 4, shown by the examples to be immunogenic. Preferably, these are used in combination with a suitable adjuvant. The isolated form of the molecule obtained via nonrecombinant means has a molecular weight of about 43 kd as determined by SDS-PAGE, and is useful in the same ways as is the recombinant protein. The recombinant form may consist of only amino acids 57-219 of the sequence of MAGE-1, as shown Also a part of the invention is the full length isolated, recombinant MAGE-1 protein, having a molecular weight of about 34.3kd as determined by SDS-PAGE, consisting of the amino acid sequence coded for by nucleotides 3931-4761 of SEQ ID NO: 1.

Other features of the invention will be clear to the artisan and need not be repeated here.

The terms and expressions which have been employed are used as terms of description and not of limitation, and there is no intention in the use of such terms and expressions of excluding any equivalents of the features shown and described or portions thereof, it being recognized that various modifications are possible within the scope of the invention.

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- (xii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Hanson, Norman D.
 - (B) REGISTRATION NUMBER: 30,946
 - (C) REFERENCE/DOCKET NUMBER: LUD 5354
- (xiii) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (212) 688-9200
 - (B) TELEFAX: (212) 838-3884

- (2) INFORMATION FOR SEQUENCE ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5674 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: genomic DNA
 - (ix) FEATURE:
 - (A) NAME/KEY: MAGE-1 gene
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

50	TCCCTCCCTT	CCACCCCAA	CCTCCCCTA	CACTGGCATC	CCCGGGGCAC
100	CAAGCCAGGC	ACCCCCAGCC	CTTCACGCTC	ATCCAAACAT	TACGCCACCC
150	AGGTGCCCAG	AGGGAAGCCC	CTCTCAACCC	TCCACCCCTG	AGAATCCGGT
200	GCGAGGTTTT	TTAGAGAGAA	GCATTAGTGG	ACTGACTTGA	ATGTGACGCC
250	CCCAGCTCTG	GGGAAGCGGG	GATCGGTGGA	GGCGGCTTGA	CGGTCTGAGG
300	CACTTACCCC	ACTGAGGACC	CTGAGGGAGG	AGGTGACATG	TAAGGAGGCA
350	ACCATCTGGT	CCAGTCCTGG	TCCCTTCATG	CCCCAAATAA	AGATAGAGGA
400	GCTTAAACCA	CCCCCTTGCT	CCACCCCAG	TCAGGCTGGG	GGTGGACTTC
450	GCTGCTTAGG	TCAGGGAAGG	CTCCGTGTGA	GAAGTCAGAG	CTGGGGACTC
500	ATTCTCAAGG	CATGCTCAGG	TGCCAGACAT	GTCCAGGCTC	AGAGGCAGC
550	CCCCACTCCA	GTGACCCAAC	CCCCACTCCC	GTCCCTAAGA	AGGGCTGAGG
600	CCAACCCCCA	CATTGTCATT	ACCCCCTCTT	CCGTGACCCA	ATGCTCACTC
650	CCGCCCAGCC	TGATGCCCAT	CCCTCAACCC	CCCACCCCAT	CCCCACATCC
700	CACCCCCACC	CGCCCACTCC	CCCACCCCA	CACCCCCACC	ATTCCACCCT
750	CGGATGTGAC	TCCGGGTGCC	CCAGGAAACA	CCGGTTCCCG	CAGGCAGGAT
800	TTCCATTCTG	GAAGCGAGGT	GGGGCAGAGA	TGCGCATTGT	GCCACTGACT
850	CTCTGTGAGG	CTGACCCAGG	CCGAAGGAAC	TAGAGTTCGG	AGGGACGGCG
900	ACTCCAAATA	GGACCCCGCC	GGAGGACTGA	AGAGGCTGAG	AGGCAAGGTG
950	CTGGCCCACC	GCTGCCAGCC	CCCCGCCCTT	AATATTCCAG	GAGAGCCCCA
1000	GCTCCAAAAG	CCAGACCCCT	TGGGCTGCCC	CGTCTCAGCC	CGCGGGAAGA
1050	CAGAGGTTGC	GCTCTGGAAT	TTCTCCCCAA	CACCAGGTTC	CCTTGAGAGA
1100	GGCTCTGCCA	GCAGGGCACA	TTAGGAGAGG	GCAGGACTGG	TGTGACCAGG
1150	CCAAGACTGC	CTGTGGGCCC	AGAGGGAGGG	TCAGCACCCA	GGCATCAAGA
1200	CCACCCAACC	TTCCCATTCC	CCCATTCGCA	CCACTCCCAC	ACTCCAATCC
1250	TACTCCGTCA	TCCCTACTCC	TCCACCCCCA	CAGCTACACC	CCCATCTCCT

CCTGACCAC	C ACCCTCCAG	C CCCAGCACCA	A GCCCCAACC	TTCTGCCACC	1300
TCACCCTCA	C TGCCCCAA	C CCCACCCTC	A TCTCTCTCAT	GTGCCCCACT	1350
CCCATCGCC	r cccccattc	r ggcagaatco	GGTTTGCCC	TGCTCTCAAC	1400
CCAGGGAAG	C CCTGGTAGG	C CCGATGTGA	ACCACTGACT	TGAACCTCAC	1450
AGATCTGAGA	A GAAGCCAGG	r TCATTTAATO	GTTCTGAGG	GCGGCTTGAG	1500
ATCCACTGAC	G GGGAGTGGT	TTAGGCTCTG	TGAGGAGGC	AGGTGAGATG	1550
CTGAGGGAGG	ACTGAGGAGG	G CACACACCC	AGGTAGATGO	CCCCAAAATG	1600
ATCCAGTACO	ACCCCTGCTC	G CCAGCCCTGG	ACCACCGGG	CAGGACAGAT	1650
GTCTCAGCTC	GACCACCCC	CGTCCCGTCC	CACTGCCACT	TAACCCACAG	1700
GGCAATCTGT	T AGTCATAGCT	TATGTGACCG	GGGCAGGGTT	GGTCAGGAGA	1750
GGCAGGGCCC	AGGCATCAAG	GTCCAGCATC	CGCCCGGCAT	TAGGGTCAGG	1800
ACCCTGGGAG	GGAACTGAGG	GTTCCCCACC	CACACCTGTC	TCCTCATCTC	1850
CACCGCCACC	CCACTCACAT	TCCCATACCT	ACCCCCTACC	CCCAACCTCA	1900
TCTTGTCAGA	ATCCCTGCTG	TCAACCCACG	GAAGCCACGG	GAATGGCGGC	1950
CAGGCACTCG	GATCTTGACG	TCCCCATCCA	GGGTCTGATG	GAGGGAAGGG	2000
GCTTGAACAG	GGCCTCAGGG	GAGCAGAGGG	AGGGCCCTAC	TGCGAGATGA	2050
GGGAGGCCTC	AGAGGACCCA	GCACCCTAGG	ACACCGCACC	CCTGTCTGAG	2100
ACTGAGGCTG	CCACTTCTGG	CCTCAAGAAT	CAGAACGATG	GGGACTCAGA	2150
TTGCATGGGG	GTGGGACCCA	GGCCTGCAAG	GCTTACGCGG	AGGAAGAGGA	2200
GGGAGGACTC	AGGGGACCTT	GGAATCCAGA	TCAGTGTGGA	CCTCGGCCCT	2250
GAGAGGTCCA	GGGCACGGTG	GCCACATATG	GCCCATATTT	CCTGCATCTT	2300
TGAGGTGACA	GGACAGAGCT	GTGGTCTGAG	AAGTGGGGCC	TCAGGTCAAC	2350
AGAGGGAGGA	GTTCCAGGAT	CCATATGGCC	CAAGATGTGC	CCCCTTCATG	2400
AGGACTGGGG	ATATCCCCGG	CTCAGAAAGA	AGGGACTCCA	CACAGTCTGG	2450
CTGTCCCCTT	TTAGTAGCTC	TAGGGGGACC	AGATCAGGGA	TGGCGGTATG	2500
TTCCATTCTC	ACTTGTACCA	CAGGCAGGAA	GTTGGGGGGC	CCTCAGGGAG	2550
ATGGGGTCTT	GGGGTAAAGG	GGGGATGTCT	ACTCATGTCA	GGGAATTGGG	2600
GGTTGAGGAA	GCACAGGCGC	TGGCAGGAAT	AAAGATGAGT	GAGACAGACA	2650
AGGCTATTGG	AATCCACACC	CCAGAACCAA	AGGGGTCAGC	CCTGGACACC	2700
rcacccagga	TGTGGCTTCT	TTTTCACTCC	TGTTTCCAGA	TCTGGGGCAG	2750
GTGAGGACCT	CATTCTCAGA	GGGTGACTCA	GGTCAACGTA	GGGACCCCCA	2800
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SAACATGAGG	GAGGACTGAG	GGTACCCCAG	GACCAGAACA	CTGAGGGAGA	2900
CTGCACAGAA	ATCAGCCCTG	CCCCTGCTGT	CACCCAGAG	AGCATGGGCT	2950
GGCCGTCTG	CCGAGGTCCT	TCCGTTATCC	TGGGATCATT	GATGTCAGGG	3000
CGGGGAGGC	CTTGGTCTGA	GAAGGCTGCG	CTCAGGTCAG	TAGAGGGAGC	3050
TCCCAGGCC	CTGCCAGGAG	TCAAGGTGAG	GACCAAGCGG	GCA CCTCA CC	2100

WO 95/20974 PCT/US95/00095

CAGG	ACAĊ	AT" T	TTAA	CCAA	T GA	ATTT	TGAT	ATC	TCTT	GCT	GCCC	TTCC	CC	3150
			CACG											3200
TCCT	TATC	AT G	GATG	TGAA	C TC	TTGA	TTTG	GAT	TTCT	CAG	ACCA	.GCAA	AA	3250
GGGC	AGGA	TC C	AGGC	CCTG	C CA	GGAA	TAAA	ATA	AGGG	ccc	TGCG	TGAG	AA	3300
CAGA	.GGGG	GT C	ATCC	ACTG	C AT	GAGA	GTGG	GGA	TGTC	ACA	GAGI	CCAG	CC	3350
CACC	CTCC	TG G	TAGC	ACTG	A GA	AGCC	AGGG	CTG	TGCT	TGC	GGTC	TGCA	CC	3400
CTGA	.GGGC	cc c	TGGA	TTCC	T CI	TCCI	'GGAG	CTC	CAGG	AAC	CAGG	CAGI	GA .	3450
GGCC	TTGG	TC I	rgaga	CAGT	A TC	CTCA	GGTC	ACA	GAGC	AGA	GGAT	GCAC	AG	3500
GGTG	TGCC	AG (CAGTG	AATG	T TI	GCCC	TGAA	TGC	ACAC	CAA	GGGC	CCCA	CC	3550
TGCC	ACAG	GA (CACAT	'AGGA	C TC	CACA	GAGI	CTG	GCCT	CAC	CTCC	CTAC	CTG	3600
TCAG	TCCI	GT A	AGAAT	CGAC	C TO	TGCI	rggcc	GGC	TGTA	ccc	TGAG	TACC	CT	3650
CTCA	CTTC	CT (CCTTC	AGGT	T TI	CAGG	GGAC	AGG	CCAA	ccc	AGAC	GAC	AGG	3700
ATTC	CCTC	GA (GCCA	CAGA	G GA	GCAC	CAAG	GAG	AAGA	TCT	GTA	AGTAC	GC	3750
CTTI	GTTA	GA (STCTC	CAAG	G TI	CAGI	TCTC	AGC	TGAG	GCC	TCT	CACAC	CAC	3800
TCCC	CTCTC	TC (CCAG	GCCI	G TO	GGT	CTTCA	A TTG	CCCA	GCT	CCT	GCCCA	ACA	3850
CTCC	TGC	TG (CTGCC	CTGA	C GA	AGAGI	CATO	2						3880
ATG	TCT	CTT	GAG	CAG	AGG	AGT	CTG	CAC	TGC	AAG	CCT	GAG	GAA	3922
GCC	CTT	GAG	GCC	CAA	CAA	GAG	GCC	CTG	GGC	CTG	GTG	TGT	GTG	3964
														4006
CTG	GAG	GAG	GTG	ccc	ACT	GCT	GGG	TCA	ACA	GAT	CCT	CCC	CAG	4048
AGT	CCT	CAG	GGA	GCC	TCC	GCC	\mathtt{TTT}	CCC	ACT	ACC	ATC	AAC	TTC	4090
ACT	CGA	CAG	AGG	CAA	ccc	AGT	GAG	GGT	TCC	AGC	AGC	CGT	GAA	4132
														4174
														4216
														4258
														4300
														4342
														4384
														4426
CTG	GGT	GAT	AAT	CAG	ATC	ATG	ccc	AAG	ACA	GGC	TTC	CTG	ATA	4468
ATT	GTC	CTG	GTC	ATG	ATT	GCA	ATG	GAG	GGC	GGC	CAT	GCT	CCT	4510
GAG	GAG	GAA	ATC	TGG	GAG	GAG	CTG	AGT	GTG	ATG	GAG	GTG	TAT	4552
GAT	GGG	AGG	GAG	CAC	AGT	GCC	TAT	GGG	GAG	CCC	AGG	AAG	CTG	4594
														4636
AGG	TGC	CGG	ACA	GTG	ATC	CCG	CAC	GCT	ATG	AGT	TCC	TGT	GGG	4678
GTC	CAA	GGG	ccc	TCG	CTG	AAA	CCA	GCT	ATG	TGA				4711
AAG'	TCCT'	TGA	GTAT	GTGA	TC A	AGGT	CAGT	G CA	AGAG	TTC				4750

	000000000			,	
			CTTTGAGAGA		4800
			AGTGGGAGGG		4850
AGTGCACCTI	CCAGGGCCGC	GTCCAGCAGC	TTCCCCTGCC	TCGTGTGACA	4900
TGAGGCCCAT	TCTTCACTCT	GAAGAGAGCG	GTCAGTGTTC	TCAGTAGTAG	4950
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			TTGAATGAAC		5050
			CTGTGTATAT		5100
			AATCCATTCT		5150
			TTAGAAATGT		5200
			AAATTAAGAG		5250
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ACCTGGATTT			TAAGAGAAAT		5350
			TCTTCTCCAT		5400
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			TCTAGGAGCT		
			GTAGGGAAAA		5500
					5550
			GTGGAGTGTC		5600
			AGTTCCTTCT		5650
GCTGGGGCAT	TTTGGGCTTT	GGGAAACTGC	AGTTCCTTCT	GGGGGAGCTG	5700
	TCTTGGGTGG				5724
					11/4

- (2) INFORMATION FOR SEQUENCE ID NO: 2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acid residues
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Ile Asn Phe Thr Arg Gln Arg Gln Pro Ser Glu Gly Ser Ser
5 10

- (2) INFORMATION FOR SEQUENCE ID NO: 3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 amino acid residues
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Leu Phe Arg Ala Val Ile Thr Lys Lys Val Ala Asp
5 10

- (2) INFORMATION FOR SEQUENCE ID NO: 4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 amino acid residues
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Asp Val Lys Glu Ala Asp Pro Thr Gly His Ser Tyr
5 10

We claim:

- Monoclonal antibody which specifically binds to tumor rejection antigen precursor MAGE-1.
- The monoclonal antibody of claim 1, designated MA454.
- 3. Hybridoma cell line which produces the monoclonal antibody of claim 1.
- 4. The hybridoma cell line of claim 3, wherein said monoclonal antibody is MA454.
- 5. Method for determining tumor rejection antigen precursor MAGE-1 in a sample, comprising contacting said sample with the monoclonal antibody of claim 1 and determining binding of said monoclonal antibody to a component of said sample as a determination of MAGE-1 in said sample.
- 6. The method of claim 5, wherein said monoclonal antibody is bound to a solid phase.
- 7. The method of claim 5, wherein said monoclonal antibody is labelled with a detectable label.
- 8. Isolated, MAGE-1 tumor rejection antigen precursor.
- 9. The isolated MAGE-1 tumor rejection antigen precursor of claim 8, which is a glycoprotein having a molecular weight of about 46 kilodaltons as determined by SDS-PAGE.
- 10. The isolated MAGE-1 tumor rejection antigen precursor of claim 8, which is a recombinantly produced protein having a molecular weight of about 34.3 kilodaltons as determined by SDS-PAGE.

- 11. Isolated protein consisting of amino acids 57-219 coded for by nucleotides 3931-4761 of the nucleotide sequence of SEQ I.D. NO.: 1.
- 12. Isolated peptide selected from the group consisting of:

SEQ ID NO: 2,

SEQ ID NO: 3, and

SEQ ID NO: 4.

- 13. Immunogenic composition comprising at least one isolated protein of claim 9 and an adjuvant.
- 14. Immunogenic composition comprising at least one isolated protein of claim 10 and an adjuvant.
- 15. Immunogenic composition comprising at least one isolated protein of claim 11 and an adjuvant.
- 16. Immunogenic composition comprising at least one peptide of claim 12 and an adjuvant.

AMENDED CLAIMS

[received by the International Bureau on 9 May 1995 (09.05.95); original claims 1-7 unchanged; original claims 8-16 replaced by new claims 8-13 (2 pages)]

- Monoclonal antibody which specifically binds to tumor rejection antigen precursor MAGE-1.
- 2. The monoclonal antibody of claim 1, designated MA454.
- 3. Hybridoma cell line which produces the monoclonal antibody of claim 1.
- 4. The hybridoma cell line of claim 3, wherein said monoclonal antibody is MA454.
- 5. Method for determining tumor rejection antigen precursor MAGE-1 in a sample, comprising contacting said sample with the monoclonal antibody of claim 1 and determining binding of said monoclonal antibody to a component of said sample as a determination of MAGE-1 in said sample.
- The method of claim 5, wherein said monoclonal antibody is bound to a solid phase.
- 7. The method of claim 5, wherein said monoclonal antibody is labelled with a detectable label.
- 8. The isolated MAGE-1 tumor rejection antigen precursor derivative which is a protein having a molecular mass of from about 20 kilodaltons to about 22 kilodaltons.
- 9. Isolated protein consisting of amino acids 57-219 coded for by nucleotides 3931-4761 of the nucleotide sequence of SEQ ID NO: 1.

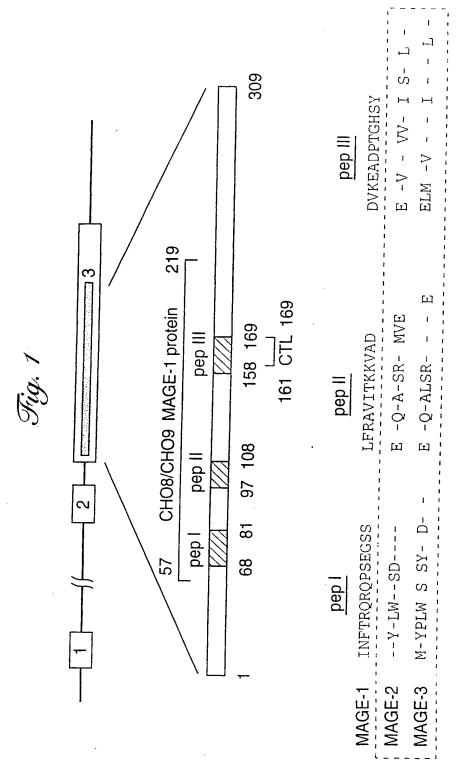
AMENDED SHEET (ARTICLE 19)

Isolated peptide selected from the group consisting 10. of:

SEQ ID NO: 2, SEQ ID NO: 3, and SEQ ID NO: 4.

- Immunogenic composition comprising at least one 11. isolated protein of claim 8 and an adjuvant.
- Immunogenic composition comprising at least one 12. isolated protein of claim 9 and an adjuvant.
- Immunogenic composition comprising at least one 13. isolated peptide of claim 10 and an adjuvant.

AMENDED SHEET (ARTICLE 19)



SUBSTITUTE SHEET (RULE 26)

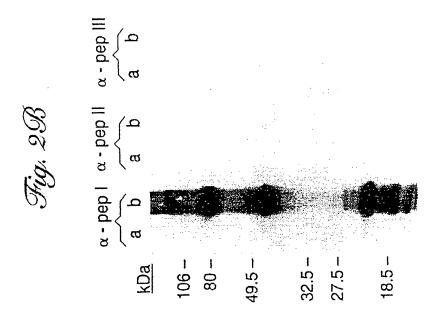
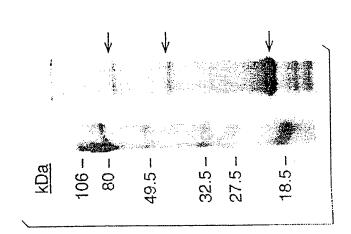
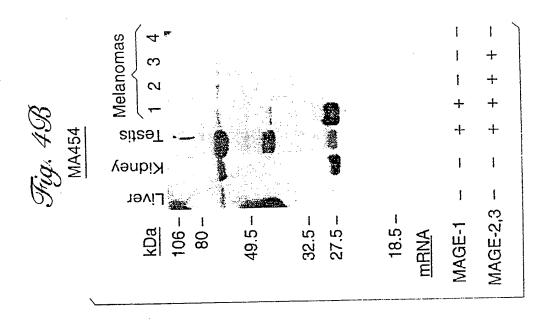


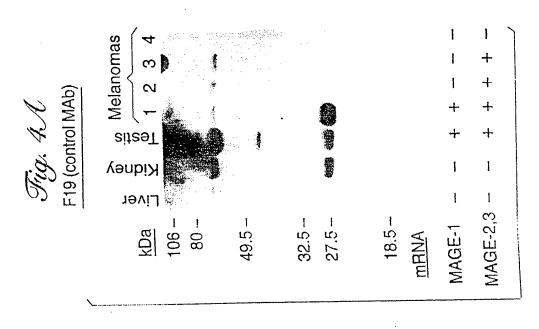
Fig. 2.A



SUBSTITUTE SHEET (RULE 26)

Fig. 36	Mel 3.1 A4554 S.2 A61 2.2 A61 2.2 ET1	-SZM	49.5 -	32.5 –	27.5 –	18.5 ~	mRNA	MAGE-1 + + -	MAGE-2,3 + + +
Fig. 3B	Rabbit anti-MAGE-1	ZZW	49.5 –	32.5 –	27.5 –	18.5 –	mRNA	MAGE-1 +	MAGE-2,3 + + -
Fig. 3M	MA454 S.2 I=M-S.2 S.2 I=M-S.2 Mel-29 -Mel-37 Mel-37	KDa NZ SK SK 106 - 80 - 80 - 80 - 80 - 80 - 80 - 80 -	TITUTE SHEET	32.5 32.5	- 5.75	18.5 –	MRNA	MAGE-1 + - + - + -	MAGE-2,3+ + + + + -



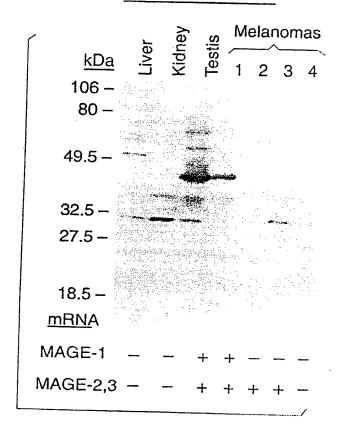


SUBSTITUTE SHEET (RULE 26)

5/5

Fig. 46

Rabbit anti-MAGE-1



SUBSTITUTE SHEET (RULE 26)

INTERNATIONAL SEARCH REPORT

International application No. PCT/US95/00095

A. CLAS	SIFICATION OF SUBJECT MATTER	•						
1PC(6) :F	1PC(6) : Please See Extra Sheet.							
US CL :F	US CL: Please See Extra Sheet. ccording to International Patent Classification (IPC) or to both national classification and IPC							
· CICLI	SC CEARCUED							
	providing searched (classification system followed by	classification symbols)	0) 2					
U.S. : 4	24/ 277.1; 435/7.1, 7.23, 240.27; 530/324, 327, 350, 3	387.7, 387.9, 388.8, 389.7, 391.1, 3	91.3					
	on searched other than minimum documentation to the ext	tent that such documents are included	in the fields searched					
	ata base consulted during the international search (name	of data base and, where practicable,	search terms used)					
	LINE, INPADOC, EMBASE, BIOSIS, REGISTRY, Herms: MAGE, antigen, peptide, protein, polypep							
search te	erms: MAGE, antigen, peptide, protein, polypep PTGHS, antibod?							
C. DOC	UMENTS CONSIDERED TO BE RELEVANT							
Category*	Citation of document, with indication, where appro	opriate, of the relevant passages	Relevant to claim No.					
X	WO, A, 92/20356 (BOON ET AL) 2		8					
 Y	entire document.		1-7, 9-11					
P, X	PROCEEDINGS OF Issued Feb	DNAL ACADEMY OF ruary 1994, Y. Chen et	1-5, 8-11					
	al., "Identification of the MAGI Monoclonal and Polyclonal Antibod see entire document.	F-I Gene House -,						
P, Y	MOLECULAR IMMUNOLOGY, Volum 1994. E. Celis et al., "Identification of of Tumor Associated Antigen MA	GE-1 for Five Common	1					
	HLA-Alleles", pages 1423-1430, se	ee entire document.						
X Fur	ther documents are listed in the continuation of Box C.	See patent family annex.						
· ;		"T" later document published after the it date and not in conflict with the appl principle or theory underlying the it	CEROOL OUT CARCA TO TITLE AND ALL AND					
•E•	to be of particular relevance carlier document published on or after the international filing date	"X" document of particular relevance; considered novel or cannot be come when the document is taken alone	the claimed invention cannot be					
	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"Y" document of particular relevance; considered to involve an invent combined with one or more other s						
.0.	document referring to an oral disclosure, use, exhibition or other means	being obvious to a person skilled in) The int					
•p•	document published prior to the international filing date but later than the priority date claimed	*& document member of the same pat						
Date of th	ne actual completion of the international search	Date of mailing of the international 03 APR 1						
27 FEB	RUARY 1995							
Commis	d mailing address of the ISA/US scioner of Patents and Trademarks	Authorized officer Line 1000 Authorized officer Anthony C. CAPUTA, Ph.	n Freder j's?					
	gton, D.C. 20231	Telephone No. (703) 308-0196						
Faccimile	No. (703) 305-3230							

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US95/00095

C (Continua	ntion). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the releva	nt passages	Relevant to claim N
Y	JOURNAL OF EXPERIMENTAL MEDICINE, Volum issued November 1992, C. Traversari et al., "A Nonap Encoded by Human Gene MAGE-1 is Recognized on H Cytolytic T Lymphocytes Directed Against Tumor Antig E", pages 1453-1457, see entire document.	eptide LA-A1 by	12-16
,	E. HARLOW et al, ANTIBODIES, A LABORATORY MANUAL", published 1988 by Cold Spring Harbor Lat (N.Y.), pages 96, 97, 139, 140, 148, 149, 553-556, and see pages 96, 97, 139, 140, 148, 149, 553-556, and 578	578-582.	1-7, 13-16
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Form PCT/ISA/210 (continuation of second sheet)(July 1992)*

INTERNATIONAL SEARCH REPORT

International application No PCT/US95/00095

	A. CLASSIFICATION OF SUBJECT MATTER: IPC (6):							
	A61K 38/04, 38/10, 38/16, 39/395, 45/00; C07K 14/46, 14/435, 16/30; C12N 5/20; G01N 33/53, 33/536							
	A. CLASSIFICATION OF SUBJECT MATTER: US CL : 424/ 277.1; 435/7.1, 7.23, 240.27; 530/324, 327, 350, 387.7, 387.9, 388.8, 389.7, 391.1, 391.3							
	424/ 277.1; 435/7.1, 7.23, 240.27; 530/324, 327, 350, 387.7, 387.9, 388.8, 389.7, 391.1, 391.3							
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